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RAW SEQUENCE LISTING   DATE: 03/27/2003
PATENT APPLICATION: US/09/889,756A   TIME: 13:19:45

        Input Set : N:\Crf4\03272003\I889756.raw
        Output Set: N:\CRF4\03272003\I889756A.raw

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 1 <110> APPLICANT: Joelle Thonnard
 2 <120> TITLE OF INVENTION: BASB055
 3 <130> FILE REFERENCE: BM45353
 4 <140> CURRENT APPLICATION NUMBER: US/09/889,756A
 5 <141> CURRENT FILING DATE: 2002-04-03
 6 <150> PRIOR APPLICATION NUMBER: 9901462.3
 7 <151> PRIOR FILING DATE: 1999-01-22
 8 <150> PRIOR APPLICATION NUMBER: 9902069.5
 9 <151> PRIOR FILING DATE: 1999-01-29
10 <160> NUMBER OF SEQ ID NOS: 4
11 <170> SOFTWARE: FastSEQ for Windows Version 4.0
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 1239
15 <212> TYPE: DNA
16 <213> ORGANISM: Neisseria meningitidis
17 <400> SEQUENCE: 1
18      atggctttt  atgctttaa  ggcgatgcgt  gcggccgcgt  tggctgccgc  cgttgcattg    60
19      gtactgtcgt  cttgcggtaa  aggcggagac  gcggcgcagg  gcgggcagcc  tgctggtcgg   120
20      gaagcccctg  cgcccgtcgt  cggtgtcgta  accgtccatc  cgcaaaccgt  cgcattgacc   180
21      gtcgagttgc  cggggcgttt  ggaatcgctg  cgtaccgccg  atgtccgcgc  ccaagtcggc   240
22      ggcatcatcc  aaaaacgcct  gttccaagaa  ggcagttatg  tccgtgccgg  acagccgctg   300
23      tatcagatcg  acagttccac  ttatgaagca  aatctggaaa  gcgcgcgcgc  gcaactggca   360
24      acggctcagg  caacgcttgc  caaagcggat  gcggatttgg  cgcgatacaa  gcctttggtt   420
25      gccgccgaag  ccgtcagccg  gcaggaatac  gatgctgcgg  taacggcgaa  acgttctgcc   480
26      gaggcaggtg  tcaaagcagc  acaggcggca  atcaaatctg  ccggcattaa  tctgaaccgt   540
27      tcgcgcatta  ccgcgccgat  ttccggcttt  atcggtcagt  ccaaagtttc  cgaaggtacg   600
28      ctgttgaatg  cgggcgatac  gaccgtgctg  gcaaccatcc  gccaaaccaa  tccgatgtat   660
29      gtgaacgtta  cccagtctgc  atccgaagtg  atgaaattgc  gccgtcagat  agccgaaggc   720
30      aaactgctgg  cggcggatgg  tgtgattgcg  gtcggcatca  aatttgacga  cggcacagtt   780
31      taccctgaaa  aaggccgcct  gctgtttgcc  gatccggtcg  tcaacgaatc  gaccggtcag   840
32      attaccctgc  gcgccgccgt  accgaacgat  cagaatatcc  tgatgcccgg  tctgtatgtg   900
33      cgcgtgctga  tggaccaagt  ggcggtggat  aacgcatttg  ttgtgccgca  gcaggcggta   960
34      acgcgcggtg  cgaaagatac  cgtgatgatt  gtgaatgccc  aaggcgggtat  ggaaccccgc  1020
35      gaggtaacgg  ttgcgcaaca  gcagggtacg  aattggattg  ttacgtcggg  tctgaaggac  1080
36      ggggacaagg  tggttgtgga  aggcatcagt  atcgccggta  taacgggtgc  gaaaaaggta  1140
37      acgcccaaag  aatgggcgtc  gtctgaaaac  caagccgccg  cgcctcaatc  cggcgttcag  1200
38      acggcatctg  aagccaaaac  tgcttctgaa  gcggaataa                           1239
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 412
42 <212> TYPE: PRT
43 <213> ORGANISM: Neisseria meningitidis
44 <400> SEQUENCE: 2
45      Met Ala Phe Tyr Ala Phe Lys Ala Met Arg Ala Ala Ala Leu Ala Ala
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```
46      1               5                  10             15
47      Ala Val Ala Leu Val Leu Ser Ser Cys Gly Lys Gly Gly Asp Ala Ala
48                  20                  25              30
49      Gln Gly Gly Gln Pro Ala Gly Arg Glu Ala Pro Ala Pro Val Val Gly
50                  35                  40              45
51      Val Val Thr Val His Pro Gln Thr Val Ala Leu Thr Val Glu Leu Pro
52          50                  55                  60
53      Gly Arg Leu Glu Ser Leu Arg Thr Ala Asp Val Arg Ala Gln Val Gly
54      65                  70                  75                  80
55      Gly Ile Ile Gln Lys Arg Leu Phe Gln Glu Gly Ser Tyr Val Arg Ala
56                  85                  90                  95
57      Gly Gln Pro Leu Tyr Gln Ile Asp Ser Ser Thr Tyr Glu Ala Asn Leu
58                  100                 105                 110
59      Glu Ser Ala Arg Ala Gln Leu Ala Thr Ala Gln Ala Thr Leu Ala Lys
60                  115                 120                 125
61      Ala Asp Ala Asp Leu Ala Arg Tyr Lys Pro Leu Val Ala Ala Glu Ala
62          130                 135                 140
63      Val Ser Arg Gln Glu Tyr Asp Ala Ala Val Thr Ala Lys Arg Ser Ala
64      145                 150                 155                 160
65      Glu Ala Gly Val Lys Ala Ala Gln Ala Ala Ile Lys Ser Ala Gly Ile
66                  165                 170                 175
67      Asn Leu Asn Arg Ser Arg Ile Thr Ala Pro Ile Ser Gly Phe Ile Gly
68                  180                 185                 190
69      Gln Ser Lys Val Ser Glu Gly Thr Leu Leu Asn Ala Gly Asp Thr Thr
70                  195                 200                 205
71      Val Leu Ala Thr Ile Arg Gln Thr Asn Pro Met Tyr Val Asn Val Thr
72          210                 215                 220
73      Gln Ser Ala Ser Glu Val Met Lys Leu Arg Arg Gln Ile Ala Glu Gly
74      225                 230                 235                 240
75      Lys Leu Leu Ala Ala Asp Gly Val Ile Ala Val Gly Ile Lys Phe Asp
76                  245                 250                 255
77      Asp Gly Thr Val Tyr Pro Glu Lys Gly Arg Leu Leu Phe Ala Asp Pro
78                  260                 265                 270
79      Val Val Asn Glu Ser Thr Gly Gln Ile Thr Leu Arg Ala Ala Val Pro
80                  275                 280                 285
81      Asn Asp Gln Asn Ile Leu Met Pro Gly Leu Tyr Val Arg Val Leu Met
82          290                 295                 300
83      Asp Gln Val Ala Val Asp Asn Ala Phe Val Val Pro Gln Gln Ala Val
84      305                 310                 315                 320
85      Thr Arg Gly Ala Lys Asp Thr Val Met Ile Val Asn Ala Gln Gly Gly
86                  325                 330                 335
87      Met Glu Pro Arg Glu Val Thr Val Ala Gln Gln Gln Gly Thr Asn Trp
88                  340                 345                 350
89      Ile Val Thr Ser Gly Leu Lys Asp Gly Asp Lys Val Val Val Glu Gly
90                  355                 360                 365
91      Ile Ser Ile Ala Gly Ile Thr Gly Ala Lys Lys Val Thr Pro Lys Glu
92          370                 375                 380
93      Trp Ala Ser Ser Glu Asn Gln Ala Ala Ala Pro Gln Ser Gly Val Gln
94      385                 390                 395                 400
```

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Output Set: **N:\CRF4\03272003\I889756A.raw**

```
 95        Thr Ala Ser Glu Ala Lys Thr Ala Ser Glu Ala Glu
 96                         405                 410
 98 <210> SEQ ID NO: 3
 99 <211> LENGTH: 41
100 <212> TYPE: DNA
101 <213> ORGANISM: Artificial Sequence
102 <220> FEATURE:
103 <223> OTHER INFORMATION: primer
104 <400> SEQUENCE: 3
105       aggcagaggc atatggcttt ttatgctttt aaggcgatgc g              41
107 <210> SEQ ID NO: 4
108 <211> LENGTH: 42
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: primer
113 <400> SEQUENCE: 4
114       aggcagaggc tcgagttccg cttcagaagc agttttggct tc             42
```

VERIFICATION SUMMARY   DATE: 03/27/2003
PATENT APPLICATION: **US/09/889,756A**   TIME: 13:19:46

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